



0118

# CRF Diskette Problem Report

The Scientific and Technical Information Center (STIC) experienced a problem when processing the following CRF diskette:

Application Serial Number:

09/036614  
3/7/98

Filing Date:

3/13/98

Classification:

Date Processed by STIC:

Point-of-Contact/Telephone No.: Arti Shah 703-308-4212

1030  
1061  
1041

1045  
04-23-98  
BUCKER'S

## Nature of Problem:

The diskette was:

- ☐ Damaged
- ☐ Unreadable
- ☐ Blank (no files present on diskette)
- ☐ Contained an empty file (filename present, but no bytes in file)
- ☐ A computer virus was detected on the diskette. The STIC will not process the diskette through the Data Capture System.

Name of virus:

The Sequence Listing was not saved in ASCII (DOS) text.

The diskette contained an error that disrupted normal processing:  
Possibly a corrupt disk sector. When STIC PC  
tried to read disk, the following message  
appeared on screen: "Data error reading drive A"  
(see attached)

SEQUENCE LISTING

09/036,614

(1) GENERAL INFORMATION

(i) APPLICANT: Hillman, Jennifer L.  
Lal, Preeti  
Tang, Y. Tom  
Yue, Henry  
Corley, Neil C.

(see last page)

(ii) TITLE OF THE INVENTION: KINESIN LIGHT CHAIN HOMOLOG

(iii) NUMBER OF SEQUENCES: 3

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Incyte Pharmaceuticals, Inc.  
(B) STREET: 3174 Porter Dr.  
(C) CITY: Palo Alto  
(D) STATE: CA  
(E) COUNTRY: USA  
(F) ZIP: 94304

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Diskette

- 09/036,614
- (B) COMPUTER: IBM Compatible
  - (C) OPERATING SYSTEM: DOS
  - (D) SOFTWARE: FastSEQ for Windows Version 2.0

(vi) CURRENT APPLICATION DATA:

- (A) APPLICATION NUMBER: To Be Assigned
- (B) FILING DATE: Filed Herewith

(vii) PRIOR APPLICATION DATA:

- (A) APPLICATION NUMBER:
- (B) FILING DATE:

(viii) ATTORNEY/AGENT INFORMATION:

- (A) NAME: Billings, Lucy J.
- (B) REGISTRATION NUMBER: 36,749
- (C) REFERENCE/DOCKET NUMBER: PF-0484 US

(ix) TELECOMMUNICATION INFORMATION:

- (A) TELEPHONE: 650-855-0555
- (B) TELEFAX: 650-845-4166

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 619 amino acids  
(B) TYPE: amino acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

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(vii) IMMEDIATE SOURCE:

(A) LIBRARY: SMCANOT01  
(B) CLONE: 2479739

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Met	Ser	Gly	Leu	Val	Leu	Gly	Gln	Arg	Asp	Glu	Pro	Ala	Gly	His	Arg
1				5				10						15	
Leu	Ser	Gln	Glu	Glu	Ile	Leu	Gly	Ser	Thr	Arg	Leu	Val	Ser	Gln	Gly
			20					25						30	
Leu	Glu	Ala	Leu	Arg	Ser	Glu	His	Gln	Ala	Val	Leu	Gln	Ser	Leu	Ser
		35					40					45			
Gln	Thr	Ile	Glu	Cys	Leu	Gln	Gln	Gly	Gly	His	Glu	Glu	Gly	Leu	Val
	50					55					60				
His	Glu	Lys	Ala	Arg	Gln	Leu	Arg	Arg	Ser	Met	Glu	Asn	Ile	Glu	Leu
65					70					75					80
Gly	Leu	Ser	Glu	Ala	Gln	Val	Met	Leu	Ala	Leu	Ala	Ser	His	Leu	Ser
			85						90					95	
Thr	Val	Glu	Ser	Glu	Lys	Gln	Lys	Leu	Arg	Ala	Gln	Val	Arg	Arg	Leu

09/036, 6/4

	MM'ÓÓÓ	19980310NE	1654	08/875,277	19980310
NE	1652	08/897,097	19980310NE	1644	08/562,983
	1654	08/977,975	19980310NE*EDIT1654	08/840,713	19980310NE
	1636	08/809,955	19980310NE*EDIT1652	08/975,522	19980310E
	1654	08/867,808	19980310NE*EDIT1654	08/902,518	A19980310NE 1
651	08/761,248	B19980310NE	1646	08/524,694	A19980310NE 163
3	08/685,589	A19980310NE*EDIT1654	08/438,762	B19980310E	1646
	08/715,202	A19980310NE	1646	08/001,773	B19980310E 1654
	08/788,943	A19980310NE*EDIT1654	08/485,218	A19980310NE	1644
08/915,498	A19980310E	1635	08/530,792	B19980310E	1652 08
/412,227	C19980310NE	1644	08/765,257	A19980310NE	1635 08/4
81,700	B19980311NE	1648	08/442,497	C19980311NE	1646 08/103
,408	A19980311NE*EDIT1634	08/661,052	19980311NE	1642	08/949,5
80	19980311NE	1649	08/833,642	19980311E	1751 08/922,315
	19980311	T	08/903,474	19980311	T 08/948,176
	19980311	T	08/871,732	B19980311E	T 08/917,456
	19980311	T	08/971,790	19980311	T 08/922,837 1
9980311	T	08/929,418	19980311	T	08/902,585 199



The above appeared in sequence I, and could be caused by a bad disk sector.

**NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING  
NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES**

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

- ☒ 1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.
- ☐ 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).
- ☐ 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
- ☐ 4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
- ☒ 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
- ☐ 6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
- ☐ 7. Other:  
**Applicant Must Provide:**
  - ☒ An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
  - ☒ An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
  - ☒ A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).

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For questions regarding compliance to these requirements, please contact:

For Rules Interpretation, call (703) 308-4216

For CRF Submission Help, call (703) 308-4212

For PatentIn software help, call (703) 308-6856

**PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE**